

## SEQUENCE LISTING

<110> Curtis, Rory A.J.

<120> 52906, 33408, AND 12189, NOVEL POTASSIUM CHANNEL FAMILY MEMBERS AND USES THEREOF

RECEIVED <130> 10448-061001 <150> US 60/209,845 JAN 3 0 2003 <151> 2000-06-06 **TECH CENTER 1600/2900** <160> 13 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 3525 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (638) ... (3178) <400> 1 60 qcqtccqcag attccagagc ctgccggctg ggaaagatcc ggtctcgggg tcggctatga 120 tecegeageg gecaaggeag ggeteaggee eegggattet eeceaeage tgetgeaetg 180 gcgcagccgg tcgccaaact ttttctcccc aaagccagtg cccccgcagt tacttggcgg qcaqccqqca gcccactctc ggcgggatga tctgggagaa gcgggcgtgg gacgaggggg 240 300 ctqctqtttt gcagccctgc gaggcgtgca gtcggagaag tggtcggggt tccacaccgt 360 ccctgagcct gcccctggc caaggtggcc cgacgtgctg cagtggctgg cgcaggtgat 420 ccqqqcaqcq cgtccggcac tagtcaaggg ggcagcggca cgggagggag gggcgccttt 480 ctettttete etececetge ageceagetg caetgegtgg gggeteteca tetecaegea 540 atcaqcaqqc ggaatccctg ccctggagcg ccctggctct ggactgcacc cccctagggt ttgtcctgca gattctcctc cccatctttc tctgccacac acgcttccct aagccgcgcg 600 655 cgccgcaaac tcagtctcgg tccccgcagg tgatgtc atg ccc att gtt ttg gtg Met Pro Ile Val Leu Val 703 cgc cca acc aat cgg act cgc cgc ctg gat tct acc gga gcc ggc atg Arg Pro Thr Asn Arg Thr Arg Arg Leu Asp Ser Thr Gly Ala Gly Met 10 20 751 ggc cct tcc tcg cac cag cag cag gag tcc ccg ctc ccg acc ata acg Gly Pro Ser Ser His Gln Gln Glu Ser Pro Leu Pro Thr Ile Thr 25 35 799 cat tgc gca ggg tgc acc acc gct tgg tct ccc tgc agc ttt aac agc His Cys Ala Gly Cys Thr Thr Ala Trp Ser Pro Cys Ser Phe Asn Ser 45 cct gac atg gaa acc cca ttg cag ttc cag cgc ggc ttc ttc cca gag 847 Pro Asp Met Glu Thr Pro Leu Gln Phe Gln Arg Gly Phe Phe Pro Glu 55 60 65

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130

135

140

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Gly Leu Val Ala Pro Gln Asn Thr Phe Leu Glu Asn Ile Val Arg Arg
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	_		_	ctc Leu										1159

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3271

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105

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Lys Asp Ile Thr Leu Phe Lys Gln Pro Ile Glu Asp Asp Ser Thr Lys 135 140

Gly Trp Thr Lys Phe Ala Arg Leu Thr Arg Ala Leu Thr Asn Ser Arg 150 155

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Lys His Ser Arg Leu Ala Glu Val Leu Gln Leu Gly Ser Asp Ile Leu 180 185 190

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His Tyr Cys Ala Phe Lys Thr Thr Trp Asp Trp Val Ile Leu Ile Leu 220 210 215

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Arg Tyr Asn Thr Ser Ala Gly Ile Trp Glu Gly Gly Pro Ser Lys Asp
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                                    410
Ser Leu Tyr Val Ser Ser Leu Tyr Phe Thr Met Thr Ser Leu Thr Thr
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Ile Gly Phe Gly Asn Ile Ala Pro Thr Thr Asp Val Glu Lys Met Phe
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Ser Val Ala Met Met Met Val Gly Ser Leu Leu Tyr Ala Thr Ile Phe
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                                            460
Gly Asn Val Thr Thr Ile Phe Gln Gln Met Tyr Ala Asn Thr Asn Arg
                                        475
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Tyr His Glu Met Leu Asn Asn Val Arg Asp Phe Leu Lys Leu Tyr Gln
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Val Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr Ile Val Ser Thr
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Trp Ser Met Ser Lys Gly Ile Asp Thr Glu Lys Val Leu Ser Ile Cys
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Pro Lys Asp Met Arg Ala Asp Ile Cys Val His Leu Asn Arg Lys Val
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Phe Asn Glu His Pro Ala Phe Arg Leu Ala Ser Asp Gly Cys Leu Arg
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Ala Leu Ala Val Glu Phe Gln Thr Ile His Cys Ala Pro Gly Asp Leu
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Ile Tyr His Ala Gly Glu Ser Val Asp Ala Leu Cys Phe Val Val Ser
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Gly Ser Leu Glu Val Ile Gln Asp Asp Glu Val Val Ala Ile Leu Gly
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Lys Gly Asp Val Phe Gly Asp Ile Phe Trp Lys Glu Thr Thr Leu Ala
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                                            620
His Ala Cys Ala Asn Val Arg Ala Leu Thr Tyr Cys Asp Leu His Ile
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                                        635
Ile Lys Arg Glu Ala Leu Leu Lys Val Leu Asp Phe Tyr Thr Ala Phe
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Ala Asn Ser Phe Ser Arg Asn Leu Thr Leu Thr Cys Asn Leu Arg Lys
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Arg Ile Ile Phe Arg Lys Ile Ser Asp Val Lys Lys Glu Glu Glu
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Arg Leu Arg Gln Lys Asn Glu Val Thr Leu Ser Ile Pro Val Asp His
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Pro Val Arg Lys Leu Phe Gln Lys Phe Lys Gln Gln Lys Glu Leu Arg
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Asn Gln Gly Ser Thr Gln Gly Asp Pro Glu Arg Asn Gln Leu Gln Val
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Glu Ser Arg Ser Leu Gln Asn Gly Thr Ser Ile Thr Gly Thr Ser Val
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Val Thr Val Ser Gln Ile Thr Pro Ile Gln Thr Ser Leu Ala Tyr Val
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Lys Thr Ser Glu Ser Leu Lys Gln Asn Asn Arg Asp Ala Met Glu Leu
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Lys Pro Asn Gly Gly Ala Asp Gln Lys Cys Leu Lys Val Asn Ser Pro
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Asn Met Gly Ala His Glu Glu Lys Lys Glu Asp Trp Asn Asn Val Thr
Lys Ala Glu Ser Met Gly Leu Leu Ser Glu Asp Pro Lys Ser Ser Asp
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                                                 845
Ser Glu Asn Ser Val Thr Lys Asn Pro Leu Arg Lys Thr Asp Ser Cys
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Asp Ser Gly Ile Thr Lys Ser Asp Leu Arg Leu Asp Lys Ala Gly Glu
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                                         875
Ala Arg Ser Pro Leu Glu His Ser Pro Ile Gln Ala Asp Ala Lys His
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                885
Pro Phe Tyr Pro Ile Pro Glu Gln Ala Leu Gln Thr Thr Leu Gln Glu
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Val Lys His Glu Leu Lys Glu Asp Ile Gln Leu Leu Ser Cys Arg Met
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Thr Ala Leu Glu Lys Gln Val Ala Glu Ile Leu Lys Ile Leu Ser Glu
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Lys Ser Val Pro Gln Ala Ser Ser Pro Lys Ser Gln Met Pro Leu Gln
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Glu Ser Pro Glu Ser Asp Lys Asp Glu Ile His Phe
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<211> 2967

<212> DNA

<213> Homo sapiens

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accaecettg eccatgeatg tgegaacgte egggeactga egtactgtga ectaeacate
                                                                      1980
atcaageggg aageettget caaagteetg gaettttata eagettttge aaaeteette
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ttacagaatg gaacctccat caccggaacc agcgtggtga ctgtgtcaca gattactccc
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atactgtcgg aaaaaagcgt accccaggcc tcatctccca aatcccaaat gccactccaa
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Cys Cys Glu Arg Leu Val Leu Asn Val Ala Gly Leu Arg Phe Glu Thr
egg geg ege egg etg gge ege tte eeg gae aet etg eta ggg gae eea
                                                                       96
Arg Ala Arg Thr Leu Gly Arg Phe Pro Asp Thr Leu Leu Gly Asp Pro
                                 25
             20
geg ege ege ege tte tae gae geg ege ege gag tat tte tte
                                                                      144
Ala Arg Arg Gly Arg Phe Tyr Asp Asp Ala Arg Arg Glu Tyr Phe Phe
         35
                             40
gac egg cae egg eee age tte gac gee gtg ete tae tae tae cag tee
                                                                      192
Asp Arg His Arg Pro Ser Phe Asp Ala Val Leu Tyr Tyr Gln Ser
     50
                         55
                                                                      240
qqt qqq cgg ctg cgg cgg ccg gcg cac gtg ccg ctc gac gtc ttc ctg
Gly Gly Arg Leu Arg Arg Pro Ala His Val Pro Leu Asp Val Phe Leu
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65				3.	70					75					80	
											gcc Ala					288
											cgc Arg					336
											gag Glu					384
	_	_		_			_	_	_		gtg Val 140	_	-			432
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_	_	_		_			_	_	_	_	gca Ala	_		_		528
	-	-	_				_		_		gga Gly					576
											acg Thr					624
											gtc Val 220					672
											gat Asp					720
											gcc Ala					768
		_	_	_	_		_	_		_	aga Arg	_		_	_	816
	-	_		_			-	_			cac His		_		_	864
											cgt Arg 300					912

ctc atc ttt ttc Leu Ile Phe Phe 305			ctc ttt tcc agc Leu Phe Ser Ser 315	
			cat ttc act agc His Phe Thr Ser	
gag tcc ttc tgg Glu Ser Phe Trp 340				
gac atg gca ccc Asp Met Ala Pro 355				
gcc att gcg ggc Ala Ile Ala Gly 370		_	cca gtg ccc gtc Pro Val Pro Val 380	
•			aca gag ggc gaa Thr Glu Gly Glu 395	
			tgt ggc cca ctg Cys Gly Pro Leu	
			gta cct gag cta Val Pro Glu Leu 430	
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tga				1341
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50	55		Leu Tyr Tyr Tyr 60	
65	70 <sup></sup>		Pro Leu Asp Val	80
Glu Glu Val Ala	Phe Tyr Gly 85	Leu Gly Ala 90	Ala Ala Leu Ala	Arg Leu 95

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Arg Glu Asp Glu Gly Cys Pro Val Pro Pro Glu Arg Pro Leu Pro Arg
                                105
            100
Arq Ala Phe Ala Arg Gln Leu Cys Leu Leu Phe Glu Phe Pro Glu Ser
                            120
Ser Gln Ala Ala Arg Val Leu Ala Val Val Ser Val Leu Val Ile Leu
                        135
                                            140
Val Ser Ile Val Val Phe Cys Leu Glu Thr Leu Pro Asp Phe Arg Asp
                                        155
                    150
Asp Arg Asp Gly Thr Gly Leu Ala Ala Ala Ala Ala Ala Gly Pro Phe
                                    170
Pro Ala Pro Leu Asn Gly Ser Ser Gln Met Pro Gly Asn Pro Pro Arg
                                185
Leu Pro Phe Asn Asp Pro Phe Phe Val Val Glu Thr Leu Cys Ile Cys
                            200
Trp Phe Ser Phe Glu Leu Leu Val Arg Leu Leu Val Cys Pro Ser Lys
                        215
                                            220
Ala Ile Phe Phe Lys Asn Val Met Asn Leu Ile Asp Phe Val Ala Ile
                                        235
                    230
Leu Pro Tyr Phe Val Ala Leu Gly Thr Glu Leu Ala Arg Gln Arg Gly
                                    250
Val Gly Gln Gln Ala Met Ser Leu Ala Ile Leu Arg Val Ile Arg Leu
                                265
            260
Val Arg Val Phe Arg Ile Phe Lys Leu Ser Arg His Ser Lys Gly Leu
                            280
Gln Ile Leu Gly Gln Thr Leu Arg Ala Ser Met Arg Glu Leu Gly Leu
                        295
                                            300
Leu Ile Phe Phe Leu Phe Ile Gly Val Val Leu Phe Ser Ser Ala Val
                    310
                                        315
Tyr Phe Ala Glu Val Asp Arg Val Asp Ser His Phe Thr Ser Ile Pro
                325
                                    330
Glu Ser Phe Trp Trp Ala Val Val Thr Met Thr Thr Val Gly Tyr Gly
                                345
            340
Asp Met Ala Pro Val Thr Val Gly Gly Lys Ile Val Gly Ser Leu Cys
                            360
Ala Ile Ala Gly Val Leu Thr Ile Ser Leu Pro Val Pro Val Ile Val
                        375
                                             380
Ser Asn Phe Ser Tyr Phe Tyr His Arg Glu Thr Glu Gly Glu Glu Ala
                    390
Gly Met Phe Ser His Val Asp Met Gln Pro Cys Gly Pro Leu Glu Gly
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                                    410
Lys Ala Asn Gly Gly Leu Val Asp Gly Glu Val Pro Glu Leu Pro Pro
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Pro Leu Trp Ala Pro Pro Gly Lys His Leu Val Thr Glu Val
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<211> 223

<212> PRT

<213> Artificial Sequence

<220>

<223> consensus sequence

<400> 9

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Ile Val Leu Lys Phe Ile Ala Tyr Gly Leu Lys Ser Thr Ser Asn Ile

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20
                                25
Ala Ala Lys Tyr Leu Lys Ser Ile Phe Asn Ile Leu Asp Leu Leu Ala
                            40
Ile Leu Pro Leu Leu Leu Leu Val Leu Phe Leu Ser Gly Thr Glu
                        55
Gln Val Ala Lys Lys Arg Leu Arg Glu Arg Phe Ser Leu Glu Leu Ser
Gln Trp Tyr Tyr Arg Ile Leu Arg Phe Leu Arg Leu Leu Arg Leu Leu
                                    90
Arq Leu Leu Arg Leu Leu Arg Leu Leu Arg Arg Leu Glu Thr Leu Phe
                                105
Glu Phe Glu Leu Gly Thr Leu Ala Trp Ser Leu Gln Ser Leu Gly Arg
                            120
Ala Leu Lys Ser Ile Leu Arg Phe Leu Leu Leu Leu Leu Leu Leu
                                            140
                        135
Ile Gly Phe Ser Val Ile Gly Tyr Leu Leu Phe Lys Gly Tyr Glu Asp
                    150
Leu Ser Glu Asn Glu Val Asp Gly Asn Ser Glu Phe Ser Ser Tyr Phe
                                    170
Asp Ala Phe Tyr Phe Leu Phe Val Thr Leu Thr Thr Val Gly Phe Gly
                                                    190
                                185
            180
Asp Leu Val Pro Val Trp Leu Gly Ile Ile Phe Phe Val Leu Phe Phe
                            200
Ile Ile Val Gly Leu Leu Leu Leu Asn Leu Leu Ile Ala Val Ile
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<213> Artificial Sequence
<220>
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Ala Leu Glu Glu Arg Ser Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln
Gly Asp Pro Gly Asp Ser Phe Tyr Ile Val Leu Ser Gly Glu Val Glu
                                25
Val Tyr Lys Leu Thr Glu Asp Gly Ala Arg Thr Pro Glu Val Ser Gln
Lys Gln Asp Thr Arg Glu Gln Val Val Ala Thr Leu Gly Pro Gly Asp
                        55
Phe Phe Gly Glu Leu Ala Leu Leu Thr Asn Asp Gly Asn Lys Asn Ala
Val Leu Pro Ser Leu Asp Gln Gly Ala Pro Arg Thr Ala Thr Val Arg
Ala Leu Thr Asp Ser Glu Leu Leu Arg Leu Asp Arg Glu Asp Phe Arg
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                                105
Arg Leu Leu Gln Lys Tyr Pro Glu
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<212> PRT

<213> Artificial Sequence

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Ser Thr Leu Thr Arg Phe Lys Pro Asp Thr Leu Leu Gly Arg Leu Leu
Lys Thr Asp Ser Asp Val His Glu Ala Arg Leu Arg Leu Cys Asp Phe
                            40
Tyr Asp Asp Glu Thr Gly Glu Tyr Phe Phe Asp Arg Ser Pro Lys His
                        55
Phe Glu Thr Ile Leu Asn Phe Tyr Arg Thr Gly Asp Gly Lys Leu His
                    70
                                        75
Arg Pro Glu Val Cys Leu Asp Ser Phe Leu Glu Glu Leu Glu Phe Tyr
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Gly Leu Asp Glu Leu Ala Ile Glu Ser Cys Cys Glu Asp Glu Tyr
                                105
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<211> 988
<212> PRT
<213> Rattus norvegicus
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Glu Asn Ile Val Arg Arg Ser Ser Glu Ser Ser Phe Leu Leu Gly Asn
Ala Gln Ile Val Asp Trp Pro Val Val Tyr Ser Asn Asp Gly Phe Cys
Lys Leu Ser Gly Tyr His Arg Ala Asp Val Met Gln Lys Ser Ser Thr
Cys Ser Phe Met Tyr Gly Glu Leu Thr Asp Lys Lys Thr Ile Glu Lys
                    70
                                         75
Val Arg Gln Thr Phe Asp Asn Tyr Glu Ser Asn Cys Phe Glu Val Leu
                                    90
Leu Tyr Lys Lys Asn Arg Thr Pro Val Trp Phe Tyr Met Gln Ile Ala
                                105
Pro Ile Arg Asn Glu His Glu Lys Val Val Leu Phe Leu Cys Thr Phe
                            120
Lys Asp Ile Thr Leu Phe Lys Gln Pro Ile Glu Asp Asp Ser Thr Lys
Gly Trp Thr Lys Phe Ala Arg Leu Thr Arg Ala Leu Thr Asn Ser Arg
                                         155
Ser Val Leu Gln Gln Leu Thr Pro Met Asn Lys Thr Glu Thr Val His
                165
                                     170
Lys His Ser Arg Leu Ala Glu Val Leu Gln Leu Gly Ser Asp Ile Leu
                                185
Pro Gln Tyr Lys Gln Glu Ala Pro Lys Thr Pro Pro His Ile Ile Leu
                            200
                                                 205
His Tyr Cys Ala Phe Lys Thr Thr Trp Asp Trp Val Ile Leu Ile Leu
                        215
                                             220
Thr Phe Tyr Thr Ala Ile Met Val Pro Tyr Asn Val Ser Phe Lys Thr
                    230
                                        235
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Lys Gln Asn Asn Ile Ala Trp Leu Val Leu Asp Ser Val Val Asp Val

250

245

Ile Phe Leu Val Asp Ile Val Leu Asn Phe His Thr Thr Phe Val Gly Pro Gly Gly Glu Val Ile Ser Asp Pro Lys Leu Ile Arg Met Asn Tyr Leu Lys Thr Trp Phe Val Ile Asp Leu Leu Ser Cys Leu Pro Tyr Asp Ile Ile Asn Ala Phe Glu Asn Val Asp Glu Gly Ile Ser Ser Leu Phe Ser Ser Leu Lys Val Val Arg Leu Leu Arg Leu Gly Arg Val Ala Arg Lys Leu Asp His Tyr Leu Glu Tyr Gly Ala Ala Val Leu Val Leu Leu Val Cys Val Phe Gly Leu Val Ala His Trp Leu Ala Cys Ile Trp Tyr Ser Ile Gly Asp Tyr Glu Val Ile Asp Glu Val Thr Asn Thr Ile Gln Ile Asp Ser Trp Leu Tyr Gln Leu Ala Leu Ser Ile Arg Thr Pro Tyr Arg Tyr Asn Thr Ser Ala Gly Ile Trp Glu Gly Gly Pro Ser Lys Asp Ser Leu Tyr Val Ser Ser Leu Tyr Phe Thr Met Thr Ser Leu Thr Thr Ile Gly Phe Gly Asn Ile Ala Pro Thr Thr Asp Val Glu Lys Met Phe Ser Val Ala Met Met Met Val Gly Ser Leu Leu Tyr Ala Thr Ile Phe Gly Asn Val Thr Thr Ile Phe Gln Gln Met Tyr Ala Asn Thr Asn Arg Tyr His Glu Met Leu Asn Asn Val Arg Asp Phe Leu Lys Leu Tyr Gln Val Pro Lys Gly Leu Ser Glu Arg Val Met Asp Tyr Ile Val Ser Thr Trp Ser Met Ser Lys Gly Ile Asp Thr Glu Lys Val Leu Ser Ile Cys Pro Lys Asp Met Arg Ala Asp Ile Cys Val His Leu Asn Arg Lys Val Phe Asn Glu His Pro Ala Phe Arg Leu Ala Ser Asp Gly Cys Leu Arg Ala Leu Ala Val Glu Phe Gln Thr Ile His Cys Ala Pro Gly Asp Leu Ile Tyr His Ala Gly Glu Ser Val Asp Ala Leu Cys Phe Val Val Ser Gly Ser Leu Glu Val Ile Gln Asp Glu Glu Val Val Ala Ile Leu Gly Lys Gly Asp Val Phe Gly Asp Ile Phe Trp Lys Glu Thr Thr Leu Ala His Ala Cys Ala Asn Val Arg Ala Leu Thr Tyr Cys Asp Leu His Ile Ile Lys Arg Glu Ala Leu Leu Lys Val Leu Asp Phe Tyr Thr Ala Phe Ala Asn Ser Phe Ser Arg Asn Leu Thr Leu Thr Cys Asn Leu Arg Lys Arg Ile Ile Phe Arg Lys Ile Ser Asp Val Lys Lys Glu Glu Glu Glu Arg Leu Arg Gln Lys Asn Glu Val Thr Leu Ser Ile Pro Val Asp His Pro Val Arg Lys Leu Phe Gln Lys Phe Lys Gln Gln Lys Glu Leu Arg

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710
Asn Gln Gly Ser Ala Gln Ser Asp Pro Glu Arg Ser Gln Leu Gln Val
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               725
Glu Ser Arg Pro Leu Gln Asn Gly Ala Ser Ile Thr Gly Thr Ser Val
                               745
Val Thr Val Ser Gln Ile Thr Pro Ile Gln Thr Ser Leu Ala Tyr Val
                           760
Lys Thr Ser Glu Thr Leu Lys Gln Asn Asn Arg Asp Ala Met Glu Leu
                       775
                                            780
Lys Pro Asn Gly Gly Ala Glu Pro Lys Cys Leu Lys Val Asn Ser Pro
                   790
                                       795
Ile Arg Met Lys Asn Gly Asn Gly Lys Gly Trp Leu Arg Leu Lys Asn
               805
                                   810
Asn Met Gly Ala His Glu Glu Lys Lys Glu Glu Trp Asn Asn Val Thr
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Lys Ala Glu Ser Met Gly Leu Leu Ser Glu Asp Pro Lys Gly Ser Asp
                            840
Ser Glu Asn Ser Val Thr Lys Asn Pro Leu Arg Lys Thr Asp Ser Cys
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Asp Ser Gly Ile Thr Lys Ser Asp Leu Arg Leu Asp Lys Ala Gly Glu
                   870
                                       875
Ala Arg Ser Pro Leu Glu His Ser Pro Ser Gln Ala Asp Ala Lys His
               885
                                   890
Pro Phe Tyr Pro Ile Pro Glu Gln Ala Leu Gln Thr Thr Leu Gln Glu
           900
                               905
Val Lys His Glu Leu Lys Glu Asp Ile Gln Leu Leu Ser Cys Arg Met
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                                               925
Thr Ala Leu Glu Lys Gln Val Ala Glu Ile Leu Lys Leu Leu Ser Glu
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                                            940
Lys Ser Val Pro Gln Thr Ser Ser Pro Lys Pro Gln Ile Pro Leu Gln
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                                        955
Val Pro Pro Gln Ile Pro Cys Gln Asp Ile Phe Ser Val Ser Arg Pro
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Glu Ser Pro Glu Ser Asp Lys Asp Glu Ile Asn Phe
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<210> 13

<211> 532

<212> PRT

<213> Mus musculus

<400> 13

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 Ala Pro Gly Gly 15

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 30

 Gly Val Thr Pro Pro Pro Pro Pro Pro Arg Pro Gly Arg Thr Phe His Ala 35
 40
 45

 Ile Phe Thr Arg Arg His Arg Thr Pro Asp Trp Gly Gly Cys Gly Val 50
 55
 60

 Gly Ala Thr Arg Pro Phe Thr Gly Arg Pro Gly Cys Ala Arg His Gly 65
 70
 75

 Ala Thr Val Pro Ala Ala Leu Arg Cys Cys Glu Arg Leu Val Leu Asn 85
 90
 95

 Val Ala Gly Leu Arg Phe Glu Thr Arg Ala Arg Thr Leu Gly Arg Phe 100
 105
 110

 Pro Asp Thr Leu Leu Gly Asp Pro Val Arg Arg Ser Arg Phe Tyr Asp
 75
 75



Gly Ala Arg Ala Glu Tyr Phe Phe Asp Arg His Arg Pro Ser Phe Asp Ala Val Leu Tyr Tyr Tyr Gln Ser Gly Gly Arg Leu Arg Arg Pro Ala His Val Pro Leu Asp Val Phe Leu Glu Glu Val Ser Phe Tyr Gly Leu Gly Arg Arg Leu Ala Arg Leu Arg Glu Asp Glu Gly Cys Ala Val Ala Glu Arg Pro Leu Pro Pro Pro Phe Ala Arg Gln Leu Trp Leu Leu Phe Glu Phe Pro Glu Ser Ser Gln Ala Ala Arg Val Leu Ala Val Val Ser Val Leu Val Ile Leu Val Ser Ile Val Val Phe Cys Leu Glu Thr Leu Pro Asp Phe Arg Asp Asp Asp Asp Pro Gly Leu Ala Pro Val Ala Ala Ala Thr Gly Ser Phe Leu Ala Arg Leu Asn Gly Ser Ser Pro Met Pro Gly Ala Pro Pro Arg Gln Pro Phe Asn Asp Pro Phe Phe Val Val Glu Thr Leu Cys Ile Cys Trp Phe Ser Phe Glu Leu Leu Val His Leu Val Ala Cys Pro Ser Lys Ala Val Phe Phe Lys Asn Val Met Asn Leu Ile Asp Phe Val Ala Ile Leu Pro Tyr Phe Val Ala Leu Gly Thr Glu Leu Ala Arg Gln Arg Gly Val Gly Gln Pro Ala Met Ser Leu Ala Ile Leu Arg Val Ile Arg Leu Val Arg Val Phe Arg Ile Phe Lys Leu Ser Arg His Ser Lys Gly Leu Gln Ile Leu Gly Gln Thr Leu Arg Ala Ser Met Arg Glu Leu Gly Leu Leu Ile Phe Phe Leu Phe Ile Gly Val Val Leu Phe Ser Ser Ala Val Tyr Phe Ala Glu Val Asp Arg Val Asp Thr His Phe Thr Ser Ile Pro Glu Ser Phe Trp Trp Ala Val Val Thr Met Thr Thr Val Gly Tyr Gly Asp Met Ala Pro Val Thr Val Gly Gly Lys Ile Val Gly Ser Leu Cys Ala Ile Ala Gly Val Leu Thr Ile Ser Leu Pro Val Pro Val Ile Val Ser Asn Phe Ser Tyr Phe Tyr His Arg Glu Thr Glu Gly Glu Glu Ala Gly Met Tyr Ser His Val Asp Thr Gln Pro Cys Gly Thr Leu Glu Gly Lys Ala Asn Gly Gly Leu Val Asp Ser Glu Val Pro Glu Leu Leu Pro Pro Leu Trp Pro Pro Ala Gly Lys His Met Val Thr Glu Val